```
                          GenCore version 6.3
                Copyright (c) 1993 - 2009  Biocceleration Ltd.


OM nucleic - nucleic search, using sw model

Run on:          June 26, 2009, 03:01:55 ; Search time 134 Seconds
                                            (without alignments)
                                            20559.273 Million cell
updates/sec

Title:           US-09-300-482-298
Perfect score:   388
Sequence:        1
ggagaaagaaagaaaagatg..........ttaaagcctgggacccccgt 388

Scoring table:   IDENTITY_NUC
                 Gapop 10.0 , Gapext 1.0

Searched:        9875436 seqs, 3558593875 residues

Total number of hits satisfying chosen parameters:      19750872

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :       Issued_Patents_NA:*
                 1:  /ABSS/Data/CRF/ptodata/1/ina/1_COMB.seq:*
                 2:  /ABSS/Data/CRF/ptodata/1/ina/5_COMB.seq:*
                 3:  /ABSS/Data/CRF/ptodata/1/ina/6A_COMB.seq:*
                 4:  /ABSS/Data/CRF/ptodata/1/ina/6B_COMB.seq:*
                 5:  /ABSS/Data/CRF/ptodata/1/ina/7A_COMB.seq:*
                 6:  /ABSS/Data/CRF/ptodata/1/ina/7B_COMB.seq:*
                 7:  /ABSS/Data/CRF/ptodata/1/ina/7C_COMB.seq:*
                 8:  /ABSS/Data/CRF/ptodata/1/ina/7D_COMB.seq:*
                 9:  /ABSS/Data/CRF/ptodata/1/ina/HA_COMB.seq:*
                 10:  /ABSS/Data/CRF/ptodata/1/ina/HB_COMB.seq:*
                 11:  /ABSS/Data/CRF/ptodata/1/ina/PCTUS_COMB.seq:*
                 12:  /ABSS/Data/CRF/ptodata/1/ina/PP_COMB.seq:*
                 13:  /ABSS/Data/CRF/ptodata/1/ina/RE_COMB.seq:*
                 14:  /ABSS/Data/CRF/ptodata/1/ina/backfiles1.seq:*
```